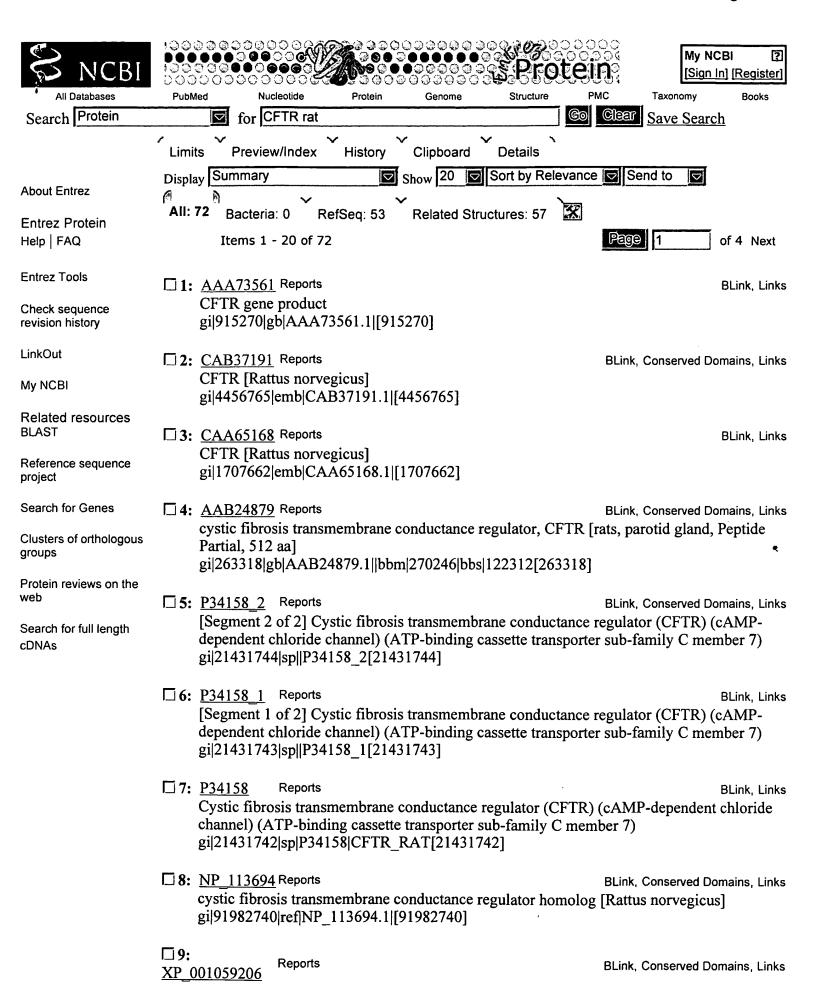


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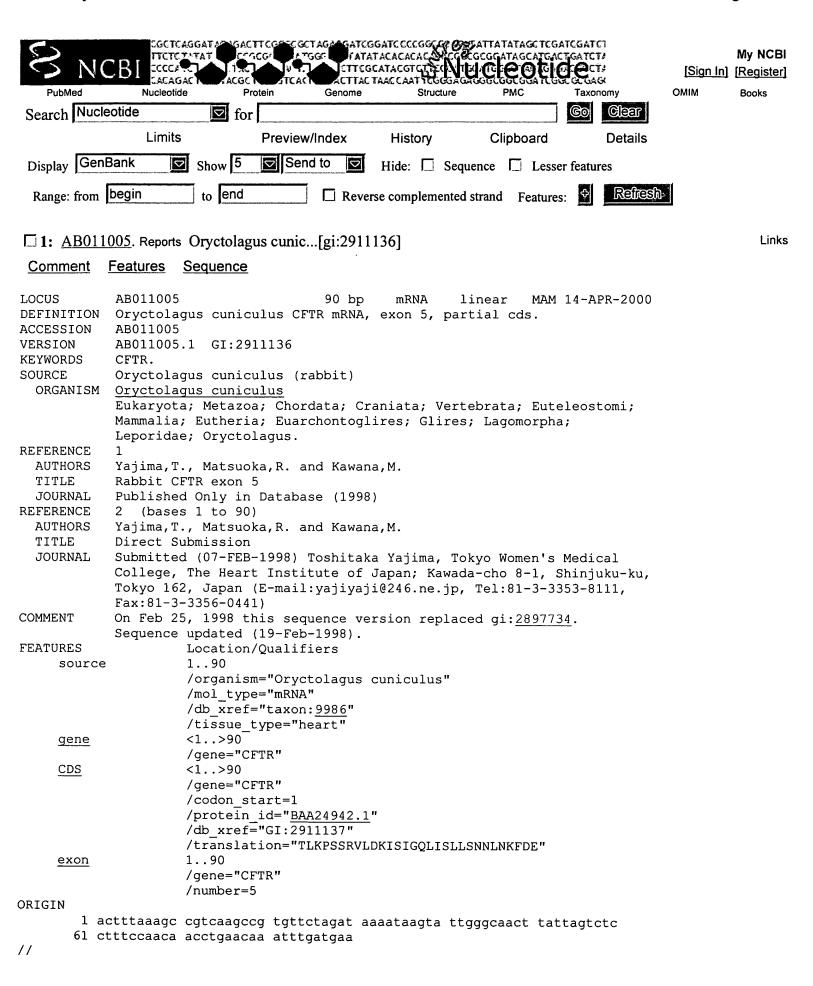


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     [Rattus norvegicus]
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□ 10:
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XP 001062374
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     [Rattus norvegicus]
     gi|109471801|ref|XP_001062374.1|[109471801]
□11:
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AAA40918
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AAR16315
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NP 066388
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□ 14: P26361
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NP 036784
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□ 16:
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NP 445876
     solute carrier family 4, sodium bicarbonate cotransporter, member 4 [Rattus norvegicus]
     gi|16758164|ref|NP 445876.1|[16758164]
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□ 18:
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NP 113736
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☐ 19: 1901178A Reports cystic fibrosis transmembrane conductance r gi 382755 prf 1901178A[382755]	BLink, Conserved Domains, Links egulator
☐ 20: Q9R1N3 Reports Sodium bicarbonate cotransporter 3 (Electro (NBC-like protein) (Solute carrier family 4 r gi 81869688 sp Q9R1N3 S4A7_RAT[81869688]	nember 7)
Items 1 - 20 of 72	Page 1 of 4 Next
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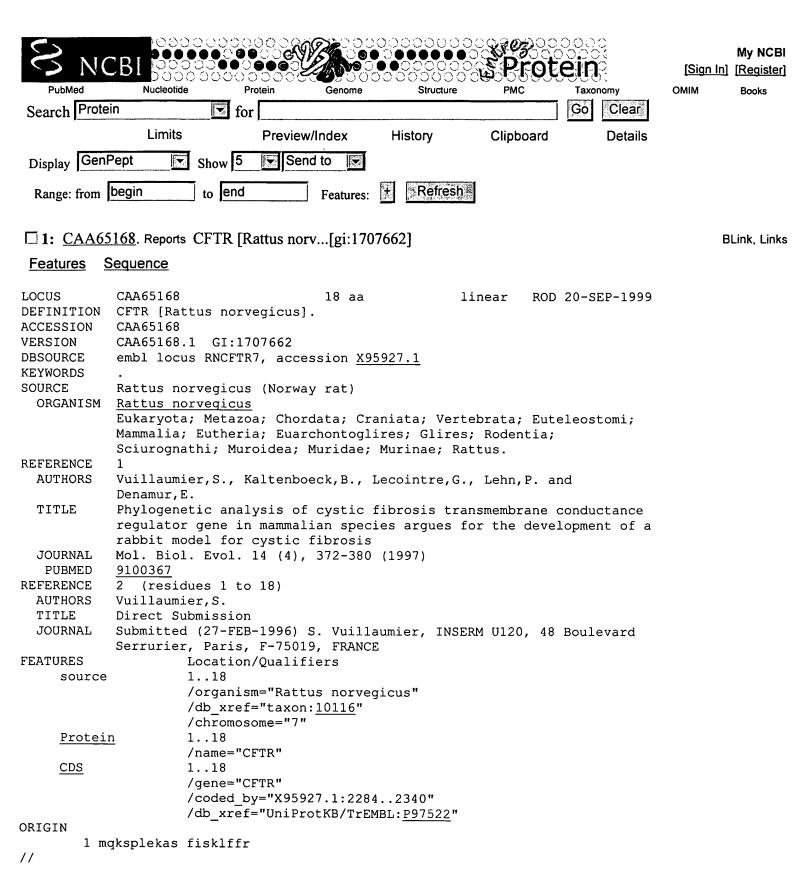
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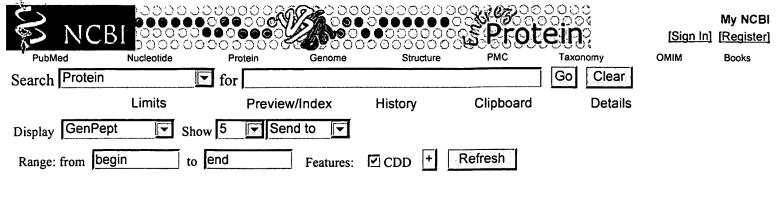


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Sep 27 2006 15:22:06



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☐ 1: <u>AAB46752</u>. Reports cystic fibrosis t...[gi:7545193]

BLink, Conserved Domains, Links

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DEFINITION
            cystic fibrosis transmembrane conductance regulator; CFTR [Mustela
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ACCESSION
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            AAB46752.2 GI:7545193
VERSION
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DBSOURCE
KEYWORDS
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            Mustela putorius furo
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Laurasiatheria; Carnivora; Caniformia;
            Mustelidae; Mustelinae; Mustela.
REFERENCE
               (residues 1 to 269)
  AUTHORS
            Sehgal, A., Presente, A. and Engelhardt, J.F.
  TITLE
            Developmental expression patterns of CFTR in ferret tracheal
            surface airway and submucosal gland epithelia
  JOURNAL
            Am. J. Respir. Cell Mol. Biol. 15 (1), 122-131 (1996)
   PUBMED
            8679216
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COMMENT
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            Method: conceptual translation supplied by author.
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translation in publication"

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L8: Entry 5 of 13

File: USPT

Mar 13, 2001

US-PAT-NO: 6201107

DOCUMENT-IDENTIFIER: US 6201107 B1

TITLE: Cystic fibrosis gene

DATE-ISSUED: March 13, 2001

INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Lap-Chee; Tsui	Toronto			CA
Riordan; John R.	Toronto			CA
Collins; Francis S.	Ann Arbor	MI		
Rommens; Johanna M.	Willowdale			CA
Iannuzzi; Michael C.	Ann Arbor	MI		
Kerem; Bat-Sheva	Toronto			CA
Drumm; Mitchell L.	Ann Arbor	MI		
Buchwald; Manuel	Toronto			CA

US-CL-CURRENT: <u>530/387.1</u>; <u>435/344</u>, <u>530/388.2</u>, <u>530/389.2</u>

CLAIMS:

What is claimed is:

- 1. An anti-CFTR polyclonal or monoclonal antibody specific for a normal CFTR polypeptide (SEQ ID NO:17), wherein said antibody is specific for an epitope of the sequence of SEQ ID NO:17 between amino acid residue positions 1 and 1480.
- 2. An anti-CFTR polyclonal ormonoclonal antibody specific for a mutant CFTR polypeptide, wherein said antibody is specific for an epitope of the sequence of SEQ ID NO:17 between amino acid residue positions 1 and 1480, wherein said amino acid sequence includes at least one cystic fibrosis (CF) mutation, wherein said cystic fibrosis (CF) mutation is a .DELTA.F508 mutation resulting from a three base pair deletion of the codon encoding phenylalanine at amino acid residue position 508 of the sequence of SEQ ID NO: 17.
- 3. A hybridoma producing a monoclonal antibody according to claim 2.
- 4. A hybridoma producing a monoclonal antibody according to claim 1.

Previous Doc Next Doc Go to Doc#

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Search Results - Record(s) 1 through 10 of 13 returned.

☐ 1. Document ID: US 6984487 B1

L8: Entry 1 of 13

File: USPT

Jan 10, 2006

US-PAT-NO: 6984487

DOCUMENT-IDENTIFIER: US 6984487 B1

TITLE: Cystic fibrosis gene

DATE-ISSUED: January 10, 2006

INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Tsui; Lap-Chee	Toronto			CA
Riordan; John R.	Toronto			CA
Collins; Francis S.	Ann Arbor	MI		US
Rommens; Johanna M.	Willowdale			CA
Iannuzzi; Michael C.	Ann Arbor	MI		US
Kerem; Bat-Sheva	Toronto			CA
Drumm; Mitchell L.	Ann Arbor	MI		US
Buchwald; Manuel	Toronto			CA

US-CL-CURRENT: 435/6; 530/350, 536/23.1

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attachments	Claims	KWIC	Draw Desc	Image
							•		•				

☐ 2. Document ID: US 6902907 B1

L8: Entry 2 of 13

File: USPT

Jun 7, 2005

US-PAT-NO: 6902907

DOCUMENT-IDENTIFIER: US 6902907 B1

TITLE: Cystic fibrosis gene

DATE-ISSUED: June 7, 2005

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY Tsui; Lap-Chee Toronto CA Riordan; John R. Toronto CA Collins; Francis S. Ann Arbor MI Rommens; Johanna M. Willowdale CA Iannuzzi; Michael C. Ann Arbor MI

Kerem; Bat-Sheva

Toronto

CA

Drumm; Mitchell L.

Ann Arbor

Buchwald; Manuel

Toronto

CA

US-CL-CURRENT: $\underline{435}/\underline{69.1}$; $\underline{435}/\underline{252.3}$, $\underline{435}/\underline{320.1}$, $\underline{435}/\underline{325}$, $\underline{530}/\underline{350}$, $\underline{536}/\underline{23.1}$

Full Titl	e Citation	Front	Review	Classification	Date	Reference	Sequences Attachmen	Claims	KMC	Draw, Desc	Image

☐ 3. Document ID: US 6730777 B1

L8: Entry 3 of 13

File: USPT

MI

May 4, 2004

US-PAT-NO: 6730777

DOCUMENT-IDENTIFIER: US 6730777 B1

TITLE: Cystic fibrosis gene

DATE-ISSUED: May 4, 2004

INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Tsui; Lap-Chee	Toronto			CA
Riordan; John R.	Toronto			CA
Collins; Francis S.	Ann Arbor	MI		
Rommens; Johanna M.	Willowdale			CA
Iannuzzi; Michael C.	Ann Arbor	MI		
Kerem; Bat-Sheva	Toronto			CA
Drumm; Mitchell L.	Ann Arbor	MI		
Buchwald; Manuel	Toronto			CA

US-CL-CURRENT: <u>530/350</u>

Full	Title	Citation	Front	Review	Classification	Date	Reference	Promines Affichiperio	Claims	KWIC	Drawu Desc	Image

☐ 4. Document ID: US 6207195 B1

L8: Entry 4 of 13

File: USPT

Mar 27, 2001

US-PAT-NO: 6207195

DOCUMENT-IDENTIFIER: US 6207195 B1

** See image for Certificate of Correction **

TITLE: Therapeutic nanospheres

DATE-ISSUED: March 27, 2001

INVENTOR-INFORMATION:

NAME CITY ZIP CODE COUNTRY STATE

Walsh; Scott Owings Mills MD Rubenstein; Ronald Baltimore MD Zeitlin; Pam Baltimore MD

Leong; Kam W.

Ellicot City

MD

US-CL-CURRENT: 424/489; 435/320.1, 435/325, 435/455, 435/458, 514/44, 977/884, 977/906, 977/915, 977/920, 977/923

Full	Title	Citation	Front	Review	Classification	Date	Reference	Sequences	Attacaments	Claims	KWIC	Draww Desc	Image

☐ 5. Document ID: US 6201107 B1

L8: Entry 5 of 13

File: USPT

Mar 13, 2001

US-PAT-NO: 6201107

DOCUMENT-IDENTIFIER: US 6201107 B1

TITLE: Cystic fibrosis gene

DATE-ISSUED: March 13, 2001

INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Lap-Chee; Tsui	Toronto			CA
Riordan; John R.	Toronto			CA
Collins; Francis S.	Ann Arbor	MI		
Rommens; Johanna M.	Willowdale			CA
Iannuzzi; Michael C.	Ann Arbor	MI		
Kerem; Bat-Sheva	Toronto			CA
Drumm; Mitchell L.	Ann Arbor	MI		
Buchwald; Manuel	Toronto			CA

US-CL-CURRENT: <u>530/387.1</u>; <u>435/344</u>, <u>530/388.2</u>, <u>530/389.2</u>

☐ 6. Document ID: US 6027880 A

L8: Entry 6 of 13

File: USPT

Feb 22, 2000

US-PAT-NO: 6027880

DOCUMENT-IDENTIFIER: US 6027880 A

TITLE: Arrays of nucleic acid probes and methods of using the same for detecting cystic

fibrosis

DATE-ISSUED: February 22, 2000

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY

Cronin; Maureen T. Los Altos CA
Miyada; Charles Garrett San Jose CA
Hubbell; Earl A. Mountain View CA
Chee; Mark Palo Alto CA

Fodor; Stephen P. A. Palo Alto CA Huang; Xiaohua C. CA Mountain View Lipshutz; Robert J. Palo Alto CA Palo Alto CA 'Lobban; Peter E. Morris; Macdonald S. Felton CA Sheldon; Edward L. San Diego CA

US-CL-CURRENT: 435/6; 422/50, 422/68.1, 436/501, 536/25.3

Full Title Citation Front Review Classification Date Reference <u>Sequences</u> Attachments Claims KMC Draw Desc Image

☐ 7. Document ID: US 6001588 A

L8: Entry 7 of 13

File: USPT

Dec 14, 1999

US-PAT-NO: 6001588

DOCUMENT-IDENTIFIER: US 6001588 A

** See image for Certificate of Correction **

TITLE: Introns and exons of the cystic fibrosis gene and mutations thereof

DATE-ISSUED: December 14, 1999

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY

Tsui; Lap-Chee Toronto CA
Rommens; Johanna M. Willowdale CA
Kerem; Bat-sheva Jerusalem IL

US-CL-CURRENT: <u>435/69.1</u>; <u>435/252.3</u>, <u>435/320.1</u>, <u>536/23.5</u>, <u>536/24.31</u>

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KWC Draw Desc Image

□ 8. Document ID: US 5981178 A

L8: Entry 8 of 13

File: USPT

Nov 9, 1999

US-PAT-NO: 5981178

DOCUMENT-IDENTIFIER: US 5981178 A

** See image for Certificate of Correction **

TITLE: Methods for screening for mutations at various positions in the introns and exons of the

cystic fibrosis gene

DATE-ISSUED: November 9, 1999

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY

Tsui; Lap-Chee Toronto CA
Rommens; Johanna M. Willowdale CA
Kerem; Bat-sheva Jerusalem IL

US-CL-CURRENT: 435/6; 435/810, 514/851, 536/24.31

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KWC Draw Desc Image

☐ 9. Document ID: US 5834421 A

L8: Entry 9 of 13

File: USPT

Nov 10, 1998

US-PAT-NO: 5834421

DOCUMENT-IDENTIFIER: US 5834421 A

TITLE: Methods and compositions for treating cystic fibrosis

DATE-ISSUED: November 10, 1998

INVENTOR-INFORMATION:

NAME

CITY

STATE

ZIP CODE

COUNTRY

Cheng; Seng Hing

Jiang; Canwen

Wellesley Marlboro MA

MA

US-CL-CURRENT: $\underline{514/2}$; $\underline{514/540}$, $\underline{514/588}$, $\underline{514/619}$, $\underline{560/33}$, $\underline{564/160}$, $\underline{564/161}$, $\underline{564/192}$, $\underline{564/59}$

Full Title Citation Front Review Classification Date Reference <u>Sequences</u> Attachments Claims KWC Draw Desc Image

☐ 10. Document ID: US 5776677 A

L8: Entry 10 of 13

File: USPT

Jul 7, 1998

US-PAT-NO: 5776677

DOCUMENT-IDENTIFIER: US 5776677 A

TITLE: Methods of detecting cystic fibrosis gene by nucleic acid hybridization

DATE-ISSUED: July 7, 1998

INVENTOR-INFORMATION:

NAME CITY STATE ZIP CODE COUNTRY Tsui; Lap-Chee Toronto CA Riordan; John R. Toronto CA Collins; Francis S. Ann Arbor MΙ Rommens; Johanna M. Willowdale CA Iannuzzi; Michael C. Ann Arbor ΜI Kerem; Bat-Sheva Toronto CA Drumm; Mitchell L. Ann Arbor ΜI Buchwald; Manuel Toronto CA

US-CL-CURRENT: 435/6; 435/91.2, 536/23.2, 536/24.3, 536/24.33

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KWC Draw Desc Image

Clear	Cenerate Collection	Polini	Fwd Reis	Blavd Refs	Generate OACS
Terms		<u> </u>		Documents	
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	L8	mutant CFTR.clm.	13
	L7	mutant CFTR	79
	L6	gene encoding cystic fibrosis transmembrane conductance regulator	39
	L5	L2 and mutation? and dna	442
	L4	L2 and mutation?	450
	L3	cystic fibrosis transmembrane conductance regulator gene.clm.	7
	L2	L1 and amino acid sequence	534
	L1	cystic fibrosis transmembrane conductance regulator	924

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=> s internalizing peptide and Mutant CFTR

L1 0 INTERNALIZING PEPTIDE AND MUTANT CFTR

=> s (CFTR or cystic fibrosis transmembrane conductance regulator) and homolog?
L2 765 (CFTR OR CYSTIC FIBROSIS TRANSMEMBRANE CONDUCTANCE REGULATOR)

AND HOMOLOG?

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=> s 14 and (mutant or mutation? variant?)

L5 44 L4 AND (MUTANT OR MUTATION? VARIANT?)

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L5 ANSWER 1 OF 44 MEDLINE on STN

ACCESSION NUMBER: 2005657059 MEDLINE

DOCUMENT NUMBER:

SOURCE:

PubMed ID: 16234241

TITLE: ATPase activity of p97/valosin-containing protein is

regulated by oxidative modification of the evolutionally

conserved cysteine 522 residue in Walker A motif.

AUTHOR: Noguchi Masakatsu; Takata Takahiro; Kimura Yoko; Manno

Atsushi; Murakami Katsuhiro; Koike Masaaki; Ohizumi

Hiroshi; Hori Seiji; Kakizuka Akira

CORPORATE SOURCE: Laboratory of Functional Biology, Kyoto University Graduate

School of Biostudies and Solution Oriented Research for

Science and Technology (JST), Kyoto 606-8501, Japan.

The Journal of biological chemistry, (2005 Dec 16) Vol.

280, No. 50, pp. 41332-41. Electronic Publication:

2005-10-18.

Journal code: 2985121R. ISSN: 0021-9258.

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 200602

ENTRY DATE: Entered STN: 18 Dec 2005

Last Updated on STN: 8 Feb 2006 Entered Medline: 7 Feb 2006

Valosin-containing protein (p97/VCP) has been proposed as playing crucial AB roles in a variety of physiological and pathological processes such as cancer and neurodegeneration. We previously showed that VCP(K524A), an ATPase activity-negative VCP mutant, induced vacuolization, accumulation of ubiquitinated proteins, and cell death, phenotypes commonly observed in neurodegenerative disorders. However, any regulatory mechanism of its ATPase activity has not yet been clarified. Here, we show that oxidative stress readily inactivates VCP ATPase activity. With liquid chromatography/tandem mass spectrometry, we found that at least three cysteine residues were modified by oxidative stress. Of them, the 522nd cysteine (Cys-522) was identified as the site responsible for the oxidative inactivation of VCP. VCP(C522T), a single-amino acid substitution mutant from cysteine to threonine, conferred almost complete resistance to the oxidative inactivation. In response to oxidative stress, VCP strengthened the interaction with Npl4 and Ufd1, both of which are essential in endoplasmic reticulum-associated protein degradation. Cys-522 is located in the second ATP binding motif and is highly conserved in multicellular but not unicellular organisms. Cdc48p (yeast VCP) has threonine in the corresponding amino acid, and it showed resistance to the oxidative inactivation in vitro. Furthermore, a yeast mutant (delta cdc48 + cdc48[T532C]) was shown to be susceptible to oxidants-induced growth inhibition and cell death. These results clearly demonstrate that VCP ATPase activity is regulated by the oxidative modification of the Cys-522 residue. This regulatory mechanism may play a key role in the conversion of oxidative stress to endoplasmic reticulum stress response in multicellular organisms and also in the pathological

L5 ANSWER 2 OF 44 MEDLINE on STN ACCESSION NUMBER: 2005088226 MEDLINE DOCUMENT NUMBER: PubMed ID: 15719171

TITLE: Binding site of activators of the cystic

process of various neurodegenerative disorders.

fibrosis transmembrane

conductance regulator in the nucleotide

binding domains.

AUTHOR: Moran O; Galietta L J V; Zegarra-Moran O

CORPORATE SOURCE: Istituto di Biofisica, CNR, Via DeMarini 6, 16149 Genoa,

Italy.. moran@ge.ibf.cnr.it

SOURCE: Cellular and molecular life sciences : CMLS, (2005 Feb)

Vol. 62, No. 4, pp. 446-60.

Journal code: 9705402. ISSN: 1420-682X.

PUB. COUNTRY: Switzerland

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 200503

ENTRY DATE: Entered STN: 19 Feb 2005

Last Updated on STN: 31 Mar 2005 Entered Medline: 30 Mar 2005

AB The use of substances that could activate the defective chloride channels of the mutant cystic fibrosis

transmembrane conductance regulator (

CFTR) has been suggested as possible therapy for cystic fibrosis.

Using epithelia formed by cells stably transfected with wildtype or mutant (G551D, G1349D) CFTR, we estimated the apparent dissociation constant, K(D), of a series of CFTR activators by measuring the increase in the apical membrane current. Modification of apparent K(D) of CFTR activators by mutations of the nucleotide-binding domains (NBDs) suggests that the binding site might be in these regions. The human NBD structure was predicted by homology with murine NBD1. An NBD1-NBD2 complex was constructed

by overlying monomers to a bacterial ABC transporter NBD dimer in the "head-to-tail" conformation. Binding sites for CFTR activators were predicted by molecular docking. Comparison of theoretical binding free energy estimated in the model to free energy estimated from the apparent dissociation constants, K(D), resulted in a remarkably good correlation coefficient for one of the putative binding sites, located in the interface between NBD1 and NBD2.

ANSWER 3 OF 44 MEDLINE on STN

ACCESSION NUMBER: 2004505132 MEDLINE DOCUMENT NUMBER: PubMed ID: 15246977

TITLE: Role of Cftr genotype in the response to chronic

Pseudomonas aeruginosa lung infection in mice.

AUTHOR: van Heeckeren Anna M; Schluchter Mark D; Drumm Mitchell L;

Davis Pamela B

CORPORATE SOURCE: Department of Pediatrics, Case Western Reserve University

School of Medicine, Cleveland, Ohio 44106-4948, USA..

anna.vanheeckeren@case.edu

CONTRACT NUMBER: HL-60293 (NHLBI)

P30 DK-27651 (NIDDK)

SOURCE: American journal of physiology. Lung cellular and molecular physiology, (2004 Nov) Vol. 287, No. 5, pp. L944-52.

Electronic Publication: 2004-07-09.

Journal code: 100901229. ISSN: 1040-0605.

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 200411

ENTRY DATE: Entered STN: 13 Oct 2004

> Last Updated on STN: 19 Dec 2004 Entered Medline: 19 Nov 2004

Patients with cystic fibrosis have a lesion in the cystic

fibrosis transmembrane conductance

regulator gene (CFTR), which is associated with abnormal regulation of other ion channels, abnormal glycosylation of secreted and cell surface molecules, and vulnerability to bacterial infection and inflammation in the lung usually leading to the death of these patients. The exact mechanism(s) by which mutation in CFTR leads to lung infection and inflammation is not clear. Mice bearing different mutations in the murine homolog to CFTR (Cftr) (R117H,

S489X, Y122X, and DeltaF508, all backcrossed to the C57BL/6J background) were compared with respect to growth and in their ability to respond to lung infection elicited with Pseudomonas aeruginosa-laden agarose beads. Body weights of mice bearing mutations in Cftr were

significantly smaller than wild-type mice at most ages. The inflammatory responses to P. aeruginosa-laden agarose beads were comparable in mice of

all four Cftr mutant genotypes with respect to absolute and relative cell counts in bronchoalveolar lavage fluid, and cytokine levels (TNF-alpha, IL-1beta, IL-6, macrophage inflammatory protein-2, and keratinocyte chemoattractant) and eicosanoid levels (PGE2 and LTB4) in epithelial lining fluid: the few small differences observed occurred only between cystic fibrosis mice bearing the S489X mutation and those bearing the knockout mutation Y122X. Thus we cannot implicate

either misprocessing of CFTR or failure of CFTR to reach the plasma membrane in the genesis of the excess inflammatory response of CF mice. Therefore, it appears that any functional defect in CFTR produces comparable inflammatory responses to lung infections with P. aeruginosa.

ANSWER 4 OF 44 MEDLINE on STN

ACCESSION NUMBER: 2004325004 MEDLINE DOCUMENT NUMBER: PubMed ID: 15225287

TITLE: Disruption of AtMRP4, a guard cell plasma membrane ABCC-type ABC transporter, leads to deregulation of stomatal opening and increased drought susceptibility.

AUTHOR: Klein Markus; Geisler Markus; Suh Su Jeoung; Kolukisaoglu H Uner; Azevedo Louis; Plaza Sonia; Curtis Mark D; Richter

Andreas; Weder Barbara; Schulz Burkhard; Martinoia Enrico

CORPORATE SOURCE: Zurich Basel Plant Science Center, University of Zurich,

Plant Biology, Zollikerstrasse 107, CH-8008 Zurich,

Switzerland.. markus.klein@botinst.unizh.ch

SOURCE: The Plant journal : for cell and molecular biology, (2004

Jul) Vol. 39, No. 2, pp. 219-36.

Journal code: 9207397. ISSN: 0960-7412.

PUB. COUNTRY: England: United Kingdom

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 200409

ENTRY DATE: Entered STN: 1 Jul 2004

Last Updated on STN: 23 Sep 2004 Entered Medline: 22 Sep 2004

AB ATP-binding cassette (ABC) transporters are membrane proteins responsible

for cellular detoxification processes in plants and animals.
Recent evidence shows that this class of transporters may also be involved

in many other cellular processes. Because of their homology with human multidrug resistance-associated proteins (MRP),

cystic fibrosis transmembrane

conductance regulator (CFTR) and sulfonylurea receptor (SUR), some plant ABC transporters have been implicated in the regulation of ion channel activities. This paper describes an investigation of the AtMRP4 gene and its role in stomatal regulation. Reporter gene studies showed that AtMRP4 is highly expressed in stomata and that the protein is localized to the plasma membrane. Stomatal aperture in three independent atmrp4 mutant alleles was larger than in wild-type plants, both in the light and in the dark, resulting in increased water loss but no change in the photosynthetic rate. In baker's yeast, AtMRP4 shows ATP-dependent, vanadate-sensitive transport of methotrevate (MTX) an antifolate and a substrate of mammalian MRPS

methotrexate (MTX), an antifolate and a substrate of mammalian MRPs. Treatment with MTX reduced stomatal opening in wild-type plants, but had no effect in atmrp4 mutants. These results indicate the involvement of AtmRP4 in the complex regulation of stomatal aperture.

L5 ANSWER 5 OF 44 MEDLINE on STN ACCESSION NUMBER: 2004131617 MEDLINE

DOCUMENT NUMBER: Publ

PubMed ID: 15024729

TITLE: Genomic rearrangements in the CFTR gene:

extensive allelic heterogeneity and diverse mutational

mechanisms.

AUTHOR:

Audrezet Marie-Pierre; Chen Jian-Min; Raguenes Odile;

Chuzhanova Nadia; Giteau Karine; Le Marechal Cedric; Quere

Isabelle; Cooper David N; Ferec Claude

CORPORATE SOURCE:

INSERM U613, Genetique Moleculaire et Genetique

Epidemiologique, Centre Hospitalier Universitaire, Brest,

France.

SOURCE:

Human mutation, (2004 Apr) Vol. 23, No. 4, pp. 343-57.

Journal code: 9215429. E-ISSN: 1098-1004.

PUB. COUNTRY: United States

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

OTHER SOURCE: OMIM-219700; OMIM-602421

ENTRY MONTH: 200405

ENTRY DATE: Entered STN: 17 Mar 2004

Last Updated on STN: 10 May 2004

Entered Medline: 6 May 2004

AB Cystic fibrosis (CF) is caused by mutations in the cystic fibrosis transmembrane conductance

regulator gene (CFTR/ABCC7). Despite the extensive and

enduring efforts of many CF researchers over the past 14 years, up to 30% of disease alleles still remain to be identified in some populations. It has long been suggested that gross genomic rearrangements could account for these unidentified alleles. To date, however, only a few large deletions have been found in the CFTR gene and only three have been fully characterized. Here, we report the first systematic screening of the 27 exons of the CFTR gene for large genomic rearrangements, by means of the quantitative multiplex PCR of short fluorescent fragments (QMPSF). A well-characterized cohort of 39 classical CF patients carrying at least one unidentified allele (after extensive and complete screening of the CFTR gene by both denaturing gradient gel electrophoresis and denaturing high-performance liquid chromatography) participated in this study. Using QMPSF, some 16% of the previously unidentified CF mutant alleles were identified and characterized, including five novel mutations (one large deletion and four indels). The breakpoints of these five mutations were precisely determined, enabling us to explore the underlying mechanisms of mutagenesis. Although non-homologous recombination may be invoked to explain all five complex lesions, each mutation appears to have arisen through a different mechanism. One of the indels was highly unusual in that it involved the insertion of a short 41 bp sequence with partial homology to a retrotranspositionally-competent LINE-1 element. The insertion of this ultra-short LINE-1 element (dubbed a "hyphen element") may constitute a novel type of mutation associated with human genetic disease.

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L5 ANSWER 6 OF 44 MEDLINE ON STN ACCESSION NUMBER: 2004003003 MEDLINE DOCUMENT NUMBER: PubMed ID: 14698290

TITLE: The rad50 signature motif: essential to ATP binding and

biological function.

AUTHOR: Moncalian Gabriel; Lengsfeld Bettina; Bhaskara Venugopal;

Hopfner Karl-Peter; Karcher Annette; Alden Erinn; Tainer

John A; Paull Tanya T

CORPORATE SOURCE: The Scripps Research Institute, 10550 North Torrey Pines

Rd., MB4, La Jolla, CA 92037, USA.

CONTRACT NUMBER: P01 CA92584 (NCI)

R01 CA94008-01 (NCI)

SOURCE: Journal of molecular biology, (2004 Jan 23) Vol. 335, No.

4, pp. 937-51.

Journal code: 2985088R. ISSN: 0022-2836.

PUB. COUNTRY: England: United Kingdom

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

OTHER SOURCE: PDB-1US8 ENTRY MONTH: 200402

ENTRY DATE: Entered STN: 6 Jan 2004

Last Updated on STN: 11 Feb 2004 Entered Medline: 10 Feb 2004

The repair of double-strand breaks in DNA is an essential process in all organisms, and requires the coordinated activities of evolutionarily conserved protein assemblies. One of the most critical of these is the Mre11/Rad50 (M/R) complex, which is present in all three biological kingdoms, but is not well-understood at the biochemical level. Previous structural analysis of a Rad50 homolog from archaebacteria illuminated the catalytic core of the enzyme, an ATP-binding domain related to the ABC transporter family of ATPases. Here, we present the crystallographic structure of the Rad50 mutant S793R. This missense signature motif mutation changes the key serine residue in the signature motif that is conserved among Rad50 homologs and ABC ATPases. The S793R mutation is analogous to the mutation S549R in the cystic fibrosis transmembrane conductance regulator (CFTR) that results in

cystic fibrosis. We show here that the serine to arginine change in the Rad50 protein prevents ATP binding and disrupts the communication among the other ATP-binding loops. This structural change, in turn, alters the communication between Rad50 monomers and thus prevents Rad50 dimerization. The equivalent mutation was made in the human Rad50 gene, and the resulting mutant protein did form a complex with Mrell and Nbs1, but was specifically deficient in all ATP-dependent enzymatic activities. This signature motif structure-function homology extends to yeast, because the same mutation introduced into the Saccharomyces cerevisiae RAD50 gene generated an allele that failed to complement a rad50 deletion strain in DNA repair assays in vivo. These structural and biochemical results extend our understanding of the Rad50 catalytic domain and validate the use of the signature motif mutant to test the role of Rad50 ATP binding in diverse organisms.

ANSWER 7 OF 44 MEDLINE on STN ACCESSION NUMBER: 2003404079 MEDLINE PubMed ID: 12907241 DOCUMENT NUMBER:

Inhibition of ATP-sensitive K+ channels by substituted TITLE:

benzo[c]quinolizinium CFTR activators.

AUTHOR: Prost Ann- Lise; Derand Renaud; Gros Laurent; Becq

Frederic; Vivaudou Michel

CEA, DRDC, Laboratoire de Biophysique Moleculaire et CORPORATE SOURCE:

Cellulaire (UMR 5090), 17 rue des Martyrs, 38054 Grenoble,

SOURCE: Biochemical pharmacology, (2003 Aug 1) Vol. 66, No. 3, pp.

425-30.

Journal code: 0101032. ISSN: 0006-2952.

PUB. COUNTRY: England: United Kingdom

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

English LANGUAGE:

Priority Journals FILE SEGMENT:

ENTRY MONTH: 200309

ENTRY DATE: Entered STN: 29 Aug 2003

Last Updated on STN: 13 Sep 2003 Entered Medline: 12 Sep 2003

AB The substituted benzo[c]quinolizinium compounds MPB-07 and MPB-91 are novel activators of the cystic fibrosis transmembrane conductance regulator (CFTR) chloride channel. High homologies between CFTR and the sulfonylurea receptor (SUR), which associates with the potassium channel Kir6.2 to form the ATP-sensitive K(+) (K(ATP)) channel, prompted us to examine possible effects of these compounds on K(ATP) channels using electrophysiological recordings and binding assays. Activity of recombinant K(ATP) channels expressed in Xenopus oocytes was recorded in the inside-out configuration of the patch-clamp technique. Channels were practically unaffected by MPB-07 but were fully blocked by MPB-91 with half-inhibition achieved at approximately 20 microM MPB-91. These effects were similar on channels formed by Kir6.2, and either the SUR1 or SUR2A isoforms were independent of the presence of nucleotides. They were not influenced by SUR mutations known to interfere with its nucleotide-binding capacity. MPB-91, but not MPB-07, was able to displace binding of glibenclamide to HEK cells expressing recombinant SUR1/Kir6.2 channels. Glibenclamide binding to native channels from pancreatic MIN6 cells was also displaced by MPB-91. A Kir6.2 mutant able to form channels without SUR was also blocked by MPB-91, but not by MPB-07. These observations demonstrate that neither MPB-07 nor MPB-91 interact with SUR, in spite of its high homology with CFTR, and that MPB-91 blocks K(ATP) channels by binding to the Kir6.2 subunit. Thus, caution should be exercised when planning to use MPB compounds in cystic fibrosis therapy, specially MPB-91 which could nonetheless find interesting applications as the precursor of a new class of K channel

blockers.

ACCESSION NUMBER: 2002423268 MEDLINE DOCUMENT NUMBER: PubMed ID: 12082160

TITLE: A role for mammalian Ubc6 homologues in

ER-associated protein degradation.

AUTHOR: Lenk Uwe; Yu Helen; Walter Jan; Gelman Marina S; Hartmann

Enno; Kopito Ron R; Sommer Thomas

CORPORATE SOURCE: The Max-Delbruck-Centrum fur Molekulare Medizin,

Robert-Rossle-Str. 10, 13092 Berlin, Germany.

SOURCE: Journal of cell science, (2002 Jul 15) Vol. 115, No. Pt 14,

pp. 3007-14.

Journal code: 0052457. ISSN: 0021-9533.

PUB. COUNTRY: England: United Kingdom

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 200212

ENTRY DATE: Entered STN: 16 Aug 2002

Last Updated on STN: 28 Dec 2002 Entered Medline: 27 Dec 2002

AB Integral membrane and secretory proteins which fail to fold productively are retained in the endoplasmic reticulum and targeted for degradation by cytoplasmic proteasomes. Genetic and biochemical analyses suggest that substrates of this pathway must be dislocated across the membrane of the endoplasmic reticulum (ER) by a process requiring a functional Sec61 complex and multiubiquitinylation. In yeast, the tail-anchored ubiquitin-conjugating enzyme Ubc6p, which is localized to the cytoplasmic surface of the ER, participates in ER-associated degradation (ERAD) of misfolded proteins. Here we describe the identification of two families of mammalian Ubc6p-related proteins. Members of both families are also located in the ER membrane and display a similar membrane topology as the yeast enzyme. Furthermore we show that expression of elevated levels of wild-type and dominant-negative alleles of these components affects specifically ERAD of the alpha subunit of the T-cell receptor and a mutant form of the CFTR protein. Similarly, we describe that the expression level of Ubc6p in yeast is also critical for ERAD, suggesting that the Ubc6p function is highly conserved from yeast to mammals.

L5 ANSWER 9 OF 44 MEDLINE on STN ACCESSION NUMBER: 2002290922 MEDLINE DOCUMENT NUMBER: PubMed ID: 12032687

TITLE: Isolation of CF cell lines corrected at DeltaF508-

CFTR locus by SFHR-mediated targeting.

AUTHOR: Bruscia E; Sangiuolo F; Sinibaldi P; Goncz K K; Novelli G;

Gruenert D C

CORPORATE SOURCE: Human Molecular Genetics, Department of Medicine,

University of Vermont, VT 05405, USA.

SOURCE: Gene therapy, (2002 Jun) Vol. 9, No. 11, pp. 683-5.

Journal code: 9421525. ISSN: 0969-7128.

PUB. COUNTRY: England: United Kingdom

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

ENTRY MONTH: 200208

ENTRY DATE: Entered STN: 29 May 2002

Last Updated on STN: 14 Aug 2002 Entered Medline: 13 Aug 2002

AB Cystic fibrosis is the most common inherited disease in the Caucasian population. About 70% of all CF chromosomes carry the DeltaF508 mutation, a 3-bp deletion that results in the loss of a phenylalanine at amino acid 508 in the CF transmembrane conductance regulator (CFTR) protein. Direct modification of the DeltaF508 locus of endogenous CFTR was achieved by small fragment homologous replacement (SFHR). Transformed human airway epithelial cells

(CFBE41o(-)), homozygous for DeltaF508 mutation, were transfected with

small fragments (491-bp) of wild-type (WT) CFTR DNA comprising exon 10 and the flanking introns. The DNA fragments were in a liposome-DNA complex at a charge ratio of 6:1 (+:-), respectively). The population of transfected cells was subcloned by limiting dilution at approximately 1 cell/well in 96-well plates. Individual colonies were isolated and analyzed. The DNA from several colonies was characterized by radiolabeled, nonallele-specific and radiolabeled, allele-specific PCR amplification, as well as by genomic DNA fingerprinting. The CFTR -WT allele was detected in five of these colonies by allele-specific PCR amplification thus indicating that the cell lines carried both WT and DeltaF alleles. DNA fingerprint analysis confirmed that the colonies were isogenic and derived from the parental CFBE41o(-) cell line. Although, the WT allele was detected by allele-specific PCR, it was not detected initially when the same samples were analyzed by non allele-specific PCR. A sensitivity assay, mixing the genomic DNA of wild-type (16HBE14o(-)) and mutant (CFBE41o(-)) cell lines, indicated that the allele-specific PCR was at least 25-fold more sensitive than non allele-specific PCR. These results suggest that the colony is not yet clonal, but still contains a population of parental, CFBE41o(-) cells that have not been modified. Based on the mixing analysis, the proportion of corrected cells appears to be between 1 and 10% of the total population. Nonallele-specific reverse transcriptase PCR (RT-PCR) analysis of the CFTR mRNA indicated that two of the colonies expressed both WT and DeltaF508 CFTR mRNA, while one colony appeared to express only the WT mRNA. The mRNA results were confirmed by sequence analysis of 3' end primer extension products from the mRNA of CFTR exon 10 showing that the mRNA containing exon 10. Furthermore, a survey of primer extension products indicated no random insertion of the fragment in an expressed gene. This study demonstrates SFHR-mediated modification of the DeltaF508 allele in DeltaF508 homozygote human airway epithelial cells over multiple generations. The resultant cells express WT-CFTR mRNA and can be subcloned further to isolate isogenic clonal populations of cells.

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FILE 'MEDLINE, HCAPLUS, BIOSIS, BIOTECHDS, EMBASE' ENTERED AT 11:48:09 ON 02 OCT 2006

- O S INTERNALIZING PEPTIDE AND MUTANT CFTR
- L2765 S (CFTR OR CYSTIC FIBROSIS TRANSMEMBRANE CONDUCTANCE REGULATOR
- L3 424 DUP REM L2 (341 DUPLICATES REMOVED)
- L4392 S L3 AND (RAT OR HUMAN OR RABBIT OR MOUSE OR ANIMAL?)
- L544 S L4 AND (MUTANT OR MUTATION? VARIANT?)
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=> s (Cystic fibrosis trans-membrane conductance regulator or CFTR) and amino acid sequenc
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L3 ANSWER 1 OF 493 HCAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER: 1998:137177 HCAPLUS

DOCUMENT NUMBER: 128:306702

TITLE: CFTR mRNA and its truncated splice variant

(TRN-CFTR) are differentially expressed

during collecting duct ontogeny

AUTHOR(S): Huber, Stephan; Braun, Gerald; Burger-Kentischer,

Anke; Reinhart, Brigitte; Luckow, Bruno; Horster,

Michael

CORPORATE SOURCE: Pettenkoferstr. 12, Physiologisches Institut,

Ludwig-Maximilians-Universitat, Munich, 80336, Germany

SOURCE: FEBS Letters (1998), 423(3), 362-366

CODEN: FEBLAL; ISSN: 0014-5793

PUBLISHER: Elsevier Science B.V.

DOCUMENT TYPE: Journal LANGUAGE: English

The collecting duct epithelium originates from the embryonic ureter by branching morphogenesis. Ontogeny-dependent changes of CFTR mRNA expression were assessed by quant. reverse transcriptase-polymerase chain reaction (RT-PCR) in primary monolayer cultures of rat ureteric buds (UB) and cortical collecting ducts, microdissected at different embryonic and postnatal developmental stages. The amt. of wild-type CFTR -specific PCR product in UB declined to 20 of the initial value between embryonic gestational day E15 and postnatal day P1. After birth the CFTR product increased transiently between P1 and P7 by a factor of 10 and decreased towards day P14. PCR products specific for TRN-CFTR, a truncated splice variant, however, were low in early embryonic cells, increased markedly between day E17 and P2, and reached a plateau postnatally. Therefore, mRNA encoding TRN-CFTR does not appear to have a specific embryonic-morphogenetic function. By contrast, such function is suggested for wild-type CFTR mRNA as its abundance was high in early embryonic nephrogenesis, as well as during a postnatal period shortly before branching morphogenesis is completed.



=> d 13 2-10

- ANSWER 2 OF 493 MEDLINE on STN L3
- AN 93122796 MEDLINE
- DN PubMed ID: 1282491
- TI Localization of the gene encoding the cystic fibrosis transmembrane conductance regulator (CFTR) in the rat to chromosome 4 and implications for the evolution of mammalian chromosomes.
- Trezise A E; Szpirer C; Buchwald M ΑU
- CS Department of Genetics, Hospital for Sick Children, Toronto, Ontario, Canada.
- so Genomics, (1992 Dec) Vol. 14, No. 4, pp. 869-74. Journal code: 8800135. ISSN: 0888-7543.

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- CY United States
- DΤ Journal; Article; (JOURNAL ARTICLE)
- LΑ English
- FS Priority Journals
- GENBANK-M89906 os
- EM 199302
- Entered STN: 26 Feb 1993 ED

Last Updated on STN: 29 Jan 1996 Entered Medline: 10 Feb 1993

- ANSWER 3 OF 493 HCAPLUS COPYRIGHT 2006 ACS on STN L3
- 2003:972178 HCAPLUS AN
- 140:35946 DN
- CFTR modifier genes and expressed proteins, in particular TΙ Kir4.2, and their regulators, useful in treating cystic fibrosis and methods and products for detecting and/or identifying same
- IN Whitsett, Jeffrey Allen; Aronow, Bruce Jefferson; Clark, Jean Cantwell
- Children's Hospital Medical Center, USA PA
- PCT Int. Appl., 80 pp. so
 - CODEN: PIXXD2
- DTPatent
- LA English
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      ANSWER 4 OF 493 HCAPLUS COPYRIGHT 2006 ACS on STN
ΑN
      2003:329918 HCAPLUS
DN
      138:380206
TI
      Alternative 5' exons of the CFTR gene show developmental
      regulation
ΑU
      Mouchel, Nathalie; Broackes-Carter, Fiona; Harris, Ann
CS
      Paediatric Molecular Genetics, Weatherall Institute of Molecular Medicine,
      John Radcliffe Hospital, Oxford University, Oxford, OX3 9DS, UK
      Human Molecular Genetics (2003), 12(7), 759-769
SO
      CODEN: HMGEE5; ISSN: 0964-6906
      Oxford University Press
PB
DТ
      Journal
     English
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               THERE ARE 24 CITED REFERENCES AVAILABLE FOR THIS RECORD
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L3
     ANSWER 5 OF 493 HCAPLUS COPYRIGHT 2006 ACS on STN
      2001:265583 HCAPLUS
AN
DN
     134:291076
TI
     Materials and method for detecting CFTR dimerization, screening
      for compounds, restoring said dimerization, as potential drugs for cystic
      fibrosis
      Teem, John L.
IN
PA
      Florida State University Research Foundation, USA
      PCT Int. Appl., 51 pp.
SO
     CODEN: PIXXD2
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     US 2000-181892P
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     ANSWER 6 OF 493 HCAPLUS COPYRIGHT 2006 ACS on STN
L3
AN
     1999:113799 HCAPLUS
DN
     130:163964
ΤI
     DNA encoding glutathione transporter function of CFTR for gene
     therapy of cystic fibrosis
     Lenoir, Gerard; Barthe, Joel; Lallemand, Jean-yves; Stoven, Veronique;
IN
     Annereau, Jean-Philippe
PA
     Assistance Publique - Hopitaux de Paris, Fr.
     PCT Int. Appl., 39 pp.
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     CODEN: PIXXD2
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                                          JP 2000-505288
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- FS Priority Journals
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- OS GENBANK-AY256886; GENBANK-AY256887; GENBANK-AY256888; GENBANK-AY256889
- EM 200407
- ED Entered STN: 13 Apr 2004 Last Updated on STN: 23 Jul 2004 Entered Medline: 22 Jul 2004

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L1 622 S (CYSTIC FIBROSIS TRANS-MEMBRANE CONDUCTANCE REGULATOR OR CFTR

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ENTRY SESSION
FULL ESTIMATED COST
DISCOUNT AMOUNTS (FOR QUALIFYING ACCOUNTS)
SINCE FILE TOTAL

CA SUBSCRIBER PRICE ENTRY SESSION -0.75 -0.75

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The vector, according to an aspect of this invention, is operatively linked to an expression control sequence in the recombinant DNA molecule so that the normal CFTR protein can be expressed, or alternatively with the other selected mutant DNA sequence the mutant CFTR polypeptide can be expressed. The expression control sequence is selected from the group consisting of sequences that control the expression of genes of prokaryotic or eukaryotic cells and their viruses and combinations thereof.

According to another aspect of the invention, a method for 10 producing normal CFTR polypeptide comprises the steps of:

- (a) culturing a host cell transfected with the recombinant vector for the normal DNA sequence in a medium and under conditions favorable for expression of the normal CFTR polypeptide; and
 - (b) isolating the expressed normal CFTR polypeptide.

According to another aspect of the invention, a method for producing a mutant CFTR polypeptide comprises the steps of:

- (a) culturing a host cell transfected with the recombinant vector for the mutant DNA sequence in a medium and under conditions favorable for expression of the mutant CFTR polypeptide; and
 - (b) isolating the expressed mutant CFTR polypeptide.

According to another aspect of the invention, a purified protein of human cell membrane origin comprises an amino sequence encoded by the mutant DNA sequence where the protein, when present in human cell membrane, is associated with cell function which causes the genetic disease cystic 30 fibrosis

According to another aspect of the invention, the CFTR polypeptide is characterized by a molecular weight of about 170,000 daltons and an epithelial cell transmembrane ion conductance affecting activity.

According to another aspect of the invention, a substantially pure CFTR protein normally expressed in human epithelial cells and characterized by being capable of participating in regulation and in control of ion transport through epithelial cells by binding to spithelial cell membrane to modulate ion movement through channels formed in the epithelial cell membrane.

According to another aspect of the invention, a process for isolating the CFTR protein comprises:

- (a) extracting peripheral proteins from membranes of epithelial cells to provide membrane material having integral proteins including said CFTR protein:
- (b) solubilizing said integral proteins of said membrane material to form a solution of said integral proteins;
- (c) separating said CFTR protein to remove any remaining other proteins of mammalian origin.

According to another aspect of the invention, a method is provided for screening a subject to determine if the subject is a CF carrier or a CF patient comprising the steps of 55 providing a biological sample of the subject to be screened and providing an assay for detecting in the biological sample, the presence of at least a member from the group consisting of the normal CF gene, normal CF gene products, a mutant CF gene, mutant CF gene products and mixtures 60 thereof.

According to another aspect of the invention, an immunologically active anti-CFTR polyclonal or monoclonal antibody specific for CFTR polypeptide is provided.

According to another aspect of the invention, a kit for 65 assaying for the presence of a CF gene by immunoassay techniques comprises:

- (a) an antibody which specifically binds to a gene product of the CF gene;
- (b) reagent means for detecting the binding of the antibody to the gene product; and
- (c) the antibody and reagent means each being present in amounts effective to perform the immunoassay.

According to another aspect of the invention, a kit for assaying for the presence of a CF gene by hybridization technique comprises:

- (a) an oligonucleotide probe which specifically binds to the CF gene;
- (b) reagent means for detecting the hybridization of the oligonucleotide probe to the CF gene; and
- (c) the probe and reagent means each being present in amounts effective to perform the hybridization assay.

According to another aspect of the invention, a method is provided for treatment for cystic fibrosis in a patient. The treatment comprises the step of administering to the patient a therapeutically effective amount of the normal CFTR protein.

According to another aspect of the invention, a method of gene therapy for cystic fibrosis comprises the step of delivery of a DNA molecule which includes a sequence corresponding to the normal DNA sequence encoding for normal CFTR protein.

According to another aspect of the invention, an animal comprises an heterologous cell system. The cell system includes a recombinant cloning vector which includes the recombinant DNA sequence corresponding to the mutant DNA sequence which induces cystic fibrosis symptoms in the animal.

According to another aspect of the invention, a transgenic mouse exhibits cystic fibrosis symptoms.

BRIEF DESCRIPTION OF THE DRAWINGS

- FIG. 1 is the nucleotide sequence of the CF gene and the amino acid sequence of the CFTR protein.
- FIG. 2 is a restriction map of the CF gene and the schematic strategy used to chromosome walk and jump to the gene.
- FIGS. 3A, 3B, 3C, 3D, and 3E are a pulse-field-gel electrophoresis map of the region including and surrounding the CF gene.
- FIGS. 4A, 4B and 4C show the detection of conserved nucleotide sequences by cross-species hybridization.
- FIG. 4D is a restriction map of overlapping segments of probes E4.3 and H1.6.
- FIG. 5 is an RNA blot hybridization analysis, using genomic and cDNA probes. Hybridization to fibroblast, trachea (normal and CF), pancreas, liver, HL60, T84, and brain RNA is shown.
- FIG. 6 is the methylation status of the E4.3 cloned region at the 5' and of the CF gene.
- FIG. 7 is a restriction map of the CFTR cDNA showing alignment of the cDNA to the genomic DNA 30 fragments.
- FIG. 8 is an RNA gel blot analysis depicting hybridization by a portion of the CFTR cDNA (clone 10-1) to a 6.5 kb mRNA transcript in various human tissues.
- FIGS. 9A, 9B, 9C, and 9D are is a DNA blot hybridization analysis depicting hybridization by the CFTR cDNA clones to genomic DNA digested with EcoRI and HindIII.
- FIGS. 10A, 10B, and 10C are is a primer extension experiment characterizing the 5' and 3' ends of the CFTR cDNA.